

SEQUENCE LISTING

<110> Fourie, Anne
Coles, Fawn
Karlsson, Lars

<120> Aggrecanase-1 and -2 Peptide Substrates and Methods

<130> ORT-1417

<160> 60

<170> PatentIn version 3.1

<210> 1
<211> 1359
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)..(1359)
<223> truncated Aggrecanase 1

<400> 1
gaattcggcca tgtcccaagac aggctcgcat cccggggagg gcttggcagg gcgctggctg 60
tggggagccc aaccctgcct cctgtcccc attgtgccgc tctcctggct ggtgtggctg 120
cttctgctac tgctggcctc tctcctgccc tcagcccgcc tggccagccc cttccccccgg 180
gaggaggaga tcgtgtttcc agagaagctc aacggcagcg tcctgcctgg ctggggcacc 240
cctgccagggc ttttgtgccc cttgcaggcc tttggggaga cgctgctact agagctggag 300
caggactccg gtgtgcaggt cgaggggctg acagtgcagt acctgggcca ggcgcctgag 360
ctgctgggtg gagcagagcc tggcacctac ctgactggca ccatcaatgg agatccggag 420
tcggtgccat ctctgcactg ggatggggaa gccctgttag gcgtgttaca atatcgggg 480
gctgaactcc acctccagcc cctggaggga ggcaccccta actctgctgg gggacctggg 540
gctcacatcc tacgcccggaa gagtcctgcc agcggtaag gtcccatgtg caacgtcaag 600
gctcctcttg gaagccccag ccccaagaccc cgaagagcca agcgctttgc ttcaactgagt 660
agatttgtgg agacactggt ggtggcagat gacaagatgg ccgcattcca cggtgccggg 720
ctaaaggcgt acctgctaac agtgatggca gcagcagcca aggccttcaa gcacccaagc 780
atccgcaatc ctgtcagctt ggtgggtgact cggctagtga tcctggggtc aggcgaggag 840
gggcccccaag tggggcccaag tgctgcccag accctgcccga gcttctgtgc ctggcagcgg 900
ggcctcaaca cccctgagga ctggaccct gaccacttg acacagccat tctgtttacc 960

cgtcaggacc	tgtgtggagt	ctccacttgc	gacacgctgg	gtatggctga	tgtgggcacc	1020
gtctgtgacc	cggctcgag	ctgtgccatt	gtggaggatg	atgggctcca	gtcagccttc	1080
actgctgctc	atgaactggg	tcatgtcttc	aacatgctcc	atgacaactc	caagccatgc	1140
atcagttga	atgggcottt	gagcacctct	cgccatgtca	tggcccctgt	gatggctcat	1200
gtggatcctg	aggagccctg	gtccccctgc	agtccccgt	tcatcactga	cttcctggac	1260
aatggctatg	ggcactgtct	cttagacaaa	ccagaggctc	cattgcacatct	gcctgtgact	1320
ggggactaca	aggacgacga	tgacaagggg	taggtcgac			1359

<210> 2
<211> 1516
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)..(1516)
<223> truncated Aggrecanse-2

<400> 2	gtcgacgcag	cgcaactatgc	tgctcggtg	ggcgtccctg	ctgctgtgcg	cgttccgcct	60
	gccccctggcc	gcgggtcggcc	ccgcccgcac	acctgcccag	gataaaagccg	ggcagcctcc	120
	gactgctgca	gcagccgccc	agcccccgcg	gcggcagggg	gaggaggtgc	aggagcgagc	180
	cgagccctccc	ggccaccgc	acccccctggc	gcagcggcgc	aggagcaagg	ggctggtgca	240
	gaacatcgac	caactcta	ccggcggcgg	caaggtggc	tacctcgtct	acgcggcgg	300
	ccgcagggttc	ctcttggacc	tggagcgaga	tggttcggtg	ggcattgctg	gcttcgtgcc	360
	cgcaggaggc	gggacgagtg	cgccctggcg	ccaccggagc	cactgcttct	atcggggcac	420
	agtggacggt	agtccccgt	ctctggctgt	ctttgacctc	tgtgggggtc	tgcacggctt	480
	cttcgcggtc	aagcacgcgc	gctacaccct	aaagccactg	ctgcgcggac	cctggggcgg	540
	ggaagaaaaag	gggcgcgtgt	acggggatgg	gtccgcacgg	atcctgcacg	tctacacccg	600
	cgagggcttc	agcttcgagg	ccctgcccgc	gcgcgcgcagc	tgcgaaaccc	ccgcgtccac	660
	accggaggcc	cacgagcatg	ctccggcgca	cagcaacccg	agcggacgcg	cagcactggc	720
	ctcgcagctc	ttggaccagt	ccgctctctc	gcccgtggg	ggctcaggac	cgcagacgtg	780
	gtggcggcgg	cggcgccgct	ccatctcccg	ggcccgccag	gtggagctgc	ttctggtgcc	840
	tgacgcgtcc	atggcgccgt	tgtatggccg	gggcctgcag	cattacctgc	tgaccctggc	900

ctccatcgcc aataggctgt acagccatgc tagcatcgag aaccacatcc gcctggccgt	960
ggtgaaggtg gtggtgctag gcgacaagga caagagcctg gaagtgagca agaacgctgc	1020
caccacactc aagaactttt gcaagtggca gcaccaacac aaccagctgg gagatgacca	1080
tgaggagcac tacgatgcag ctatcctgtt tactcggag gatttatgtg ggcatttcattc	1140
atgtgacacc ctgggaatgg cagacgttgg gaccatatgt tctccagagc gcagctgtgc	1200
tgtgattgaa gacgatggcc tccacgcagc cttcaactgtg gtcacgaaa tcggacattt	1260
acttggcctc tccccatgacg attccaaatt ctgtgaagag acctttggtt ccacagaaga	1320
taagcgctta atgtcttcca tccttaccag cattgatgca tctaagccct ggtccaaatg	1380
cacttcagcc accatcacag aattcctgga tgatggccat ggtaactgtt tgctggacct	1440
accacgaaag cagatcctgg gcggggacta caaggacgac gatgacaagg ggtagaagct	1500
tqtcqaqaq tactaq	1516

<210> 3
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> peptide substrate

<400> 3

Lys Glu Leu Ala Glu Leu Arg Glu Ser Thr Ser
 1 5 10

```
<210> 4
<211> 11
<212> PRT
<213> Artificial Sequence
```

<220>
<223> Peptide substrate

<400> 4

Ala Asp Leu Ser Ser Phe Lys Ser Gln Glu Leu
1 5 10

```
<210> 5
<211> 10
<212> PRT
<213> Artificial sequence
```

<220>
<223> Peptide substrate

<400> 5

Glu Lys Ala Arg Val Leu Ala Glu Ala Ala
1 5 10

<210> 6
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide Substrate

<400> 6

Glu Lys Ala Arg Val Leu Ala Glu Ala Met
1 5 10

<210> 7
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide substrate

<400> 7

Glu Arg Ala Glu Gln Gln Arg Leu Lys Ser Gln Asp Leu
1 5 10

<210> 8
<211> 447
<212> PRT
<213> Homo sapiens

<220>
<221> MISC_FEATURE
<222> (1)..(447)
<223> truncated Aggrecanase 1

<400> 8

Met Ser Gln Thr Gly Ser His Pro Gly Arg Gly Leu Ala Gly Arg Trp
1 5 10 15

Leu Trp Gly Ala Gln Pro Cys Leu Leu Leu Pro Ile Val Pro Leu Ser
20 25 30

Trp Leu Val Trp Leu Leu Leu Leu Ala Ser Leu Leu Pro Ser
 35 40 45

Ala Arg Leu Ala Ser Pro Leu Pro Arg Glu Glu Glu Ile Val Phe Pro
 50 55 60

Glu Lys Leu Asn Gly Ser Val Leu Pro Gly Ser Gly Thr Pro Ala Arg
 65 70 75 80

Leu Leu Cys Arg Leu Gln Ala Phe Gly Glu Thr Leu Leu Leu Glu Leu
 85 90 95

Glu Gln Asp Ser Gly Val Gln Val Glu Gly Leu Thr Val Gln Tyr Leu
 100 105 110

Gly Gln Ala Pro Glu Leu Leu Gly Gly Ala Glu Pro Gly Thr Tyr Leu
 115 120 125

Thr Gly Thr Ile Asn Gly Asp Pro Glu Ser Val Ala Ser Leu His Trp
 130 135 140

Asp Gly Gly Ala Leu Leu Gly Val Leu Gln Tyr Arg Gly Ala Glu Leu
 145 150 155 160

His Leu Gln Pro Leu Glu Gly Gly Thr Pro Asn Ser Ala Gly Gly Pro
 165 170 175

Gly Ala His Ile Leu Arg Arg Lys Ser Pro Ala Ser Gly Gln Gly Pro
 180 185 190

Met Cys Asn Val Lys Ala Pro Leu Gly Ser Pro Ser Pro Arg Pro Arg
 195 200 205

Arg Ala Lys Arg Phe Ala Ser Leu Ser Arg Phe Val Glu Thr Leu Val
 210 215 220

Val Ala Asp Asp Lys Met Ala Ala Phe His Gly Ala Gly Leu Lys Arg
 225 230 235 240

Tyr Leu Leu Thr Val Met Ala Ala Ala Lys Ala Phe Lys His Pro
 245 250 255

Ser Ile Arg Asn Pro Val Ser Leu Val Val Thr Arg Leu Val Ile Leu
 260 265 270

Gly Ser Gly Glu Glu Gly Pro Gln Val Gly Pro Ser Ala Ala Gln Thr
 275 280 285

Leu Arg Ser Phe Cys Ala Trp Gln Arg Gly Leu Asn Thr Pro Glu Asp
 290 295 300

Ser Asp Pro Asp His Phe Asp Thr Ala Ile Leu Phe Thr Arg Gln Asp
 305 310 315 320

Leu Cys Gly Val Ser Thr Cys Asp Thr Leu Gly Met Ala Asp Val Gly
 325 330 335

Thr Val Cys Asp Pro Ala Arg Ser Cys Ala Ile Val Glu Asp Asp Gly
 340 345 350

Leu Gln Ser Ala Phe Thr Ala Ala His Glu Leu Gly His Val Phe Asn
 355 360 365

Met Leu His Asp Asn Ser Lys Pro Cys Ile Ser Leu Asn Gly Pro Leu
 370 375 380

Ser Thr Ser Arg His Val Met Ala Pro Val Met Ala His Val Asp Pro
 385 390 395 400

Glu Glu Pro Trp Ser Pro Cys Ser Ala Arg Phe Ile Thr Asp Phe Leu
 405 410 415

Asp Asn Gly Tyr Gly His Cys Leu Leu Asp Lys Pro Glu Ala Pro Leu
 420 425 430

His Leu Pro Val Thr Gly Asp Tyr Lys Asp Asp Asp Lys Gly
 435 440 445

<210> 9
 <211> 492
 <212> PRT
 <213> Homo sapiens

<220>
 <221> MISC_FEATURE
 <222> (1)..(492)
 <223> truncated Aggrecanse-2

<400> 9

Met Leu Leu Gly Trp Ala Ser Leu Leu Leu Cys Ala Phe Arg Leu Pro
 1 5 10 15

Leu Ala Ala Val Gly Pro Ala Ala Thr Pro Ala Gln Asp Lys Ala Gly
 20 25 30

Gln Pro Pro Thr Ala Ala Ala Ala Gln Pro Arg Arg Arg Gln Gly
 35 40 45

Glu Glu Val Gln Glu Arg Ala Glu Pro Pro Gly His Pro His Pro Leu
 50 55 60

Ala Gln Arg Arg Arg Ser Lys Gly Leu Val Gln Asn Ile Asp Gln Leu
 65 70 75 80

Tyr Ser Gly Gly Lys Val Gly Tyr Leu Val Tyr Ala Gly Gly Arg
 85 90 95

Arg Phe Leu Leu Asp Leu Glu Arg Asp Gly Ser Val Gly Ile Ala Gly
 100 105 110

Phe Val Pro Ala Gly Gly Thr Ser Ala Pro Trp Arg His Arg Ser
 115 120 125

His Cys Phe Tyr Arg Gly Thr Val Asp Gly Ser Pro Arg Ser Leu Ala
 130 135 140

Val Phe Asp Leu Cys Gly Gly Leu Asp Gly Phe Phe Ala Val Lys His
 145 150 155 160

Ala Arg Tyr Thr Leu Lys Pro Leu Leu Arg Gly Pro Trp Ala Glu Glu
 165 170 175

Glu Lys Gly Arg Val Tyr Gly Asp Gly Ser Ala Arg Ile Leu His Val
 180 185 190

Tyr Thr Arg Glu Gly Phe Ser Phe Glu Ala Leu Pro Pro Arg Ala Ser
 195 200 205

Cys Glu Thr Pro Ala Ser Thr Pro Glu Ala His Glu His Ala Pro Ala

210

215

220

His Ser Asn Pro Ser Gly Arg Ala Ala Leu Ala Ser Gln Leu Leu Asp
 225 230 235 240

Gln Ser Ala Leu Ser Pro Ala Gly Gly Ser Gly Pro Gln Thr Trp Trp
 245 250 255

Arg Arg Arg Arg Ser Ile Ser Arg Ala Arg Gln Val Glu Leu Leu
 260 265 270

Leu Val Ala Asp Ala Ser Met Ala Arg Leu Tyr Gly Arg Gly Leu Gln
 275 280 285

His Tyr Leu Leu Thr Leu Ala Ser Ile Ala Asn Arg Leu Tyr Ser His
 290 295 300

Ala Ser Ile Glu Asn His Ile Arg Leu Ala Val Val Lys Val Val Val
 305 310 315 320

Leu Gly Asp Lys Asp Lys Ser Leu Glu Val Ser Lys Asn Ala Ala Thr
 325 330 335

Thr Leu Lys Asn Phe Cys Lys Trp Gln His Gln His Asn Gln Leu Gly
 340 345 350

Asp Asp His Glu Glu His Tyr Asp Ala Ala Ile Leu Phe Thr Arg Glu
 355 360 365

Asp Leu Cys Gly His His Ser Cys Asp Thr Leu Gly Met Ala Asp Val
 370 375 380

Gly Thr Ile Cys Ser Pro Glu Arg Ser Cys Ala Val Ile Glu Asp Asp
 385 390 395 400

Gly Leu His Ala Ala Phe Thr Val Ala His Glu Ile Gly His Leu Leu
 405 410 415

Gly Leu Ser His Asp Asp Ser Lys Phe Cys Glu Glu Thr Phe Gly Ser
 420 425 430

Thr Glu Asp Lys Arg Leu Met Ser Ser Ile Leu Thr Ser Ile Asp Ala
 435 440 445

Ser Lys Pro Trp Ser Lys Cys Thr Ser Ala Thr Ile Thr Glu Phe Leu
450 455 460

Asp Asp Gly His Gly Asn Cys Leu Leu Asp Leu Pro Arg Lys Gln Ile
465 470 475 480

Leu Gly Gly Asp Tyr Lys Asp Asp Asp Lys Gly
485 490

<210> 10
<211> 13
<212> PRT
<213> Artificial sequence

<220>
<223> Peptide substrate

<400> 10

Glu His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg
1 5 10

<210> 11
<211> 4
<212> PRT
<213> Artificial sequence

<220>
<223> Peptide substrate

<400> 11

Glu Ala Glu Asn
1

<210> 12
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide Substrate

<400> 12

Glu Gly Arg His Ile Asp Asn Glu Glu Asp Ile
1 5 10

<210> 13

<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide substrate

<400> 13

Glu Gly Asn Ala Phe Asn Asn Leu Asp
1 5

<210> 14
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide substrate

<400> 14

Glu Tyr Thr Pro Asn Asn Glu Ile Asp Ser Phe
1 5 10

<210> 15
<211> 8
<212> PRT
<213> Artificial sequence

<220>
<223> Peptide substrate

<400> 15

Glu Gln Leu Arg Met Lys Leu Pro
1 5

<210> 16
<211> 8
<212> PRT
<213> Artificial sequence

<220>
<223> Peptide substrate

<400> 16

Glu Arg Gly Phe Phe Tyr Thr Pro
1 5

<210> 17
<211> 8

<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide substrate

<400> 17

Glu Val Thr Glu Gly Pro Ile Pro
1 5

<210> 18
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide substrate

<400> 18

Glu Pro Leu Phe Tyr Glu Ala Pro
1 5

<210> 19
<211> 8
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 19

Glu Leu Pro Met Gly Ala Leu Pro
1 5

<210> 20
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide substrate

<400> 20

Glu Lys Pro Ala Ala Phe Phe Arg Leu
1 5

<210> 21
<211> 13
<212> PRT

<213> Artificial sequence

<220>

<223> peptide substrate .

<400> 21

Glu Leu Tyr Glu Asn Lys Pro Arg Arg Pro Tyr Ile Leu
1 5 10

<210> 22

<211> 10

<212> PRT

<213> Artificial sequence

<220>

<223> peptide substrate

<400> 22

Glu Ser Glu Val Asn Leu Asp Ala Glu Phe
1 5 10

<210> 23

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide Substrate

<400> 23

Glu Ser Gln Asn Tyr Pro Ile Val Gln
1 5

<210> 24

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide Substrate

<400> 24

Glu Lys Pro Ile Glu Phe Phe Arg Leu
1 5

<210> 25

<211> 9

<212> PRT

<213> Artificial sequence

<220>
<223> Peptide substrate

<400> 25

Glu Lys Pro Ala Glu Phe Phe Ala Leu
1 5

<210> 26
<211> 9
<212> PRT
<213> Artificial sequence

<220>
<223> peptide substrate

<400> 26

Glu Lys Pro Ala Lys Phe Phe Arg Leu
1 5

<210> 27
<211> 10
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 27

Glu Ile Pro Phe His Leu Val Ile His Thr
1 5 10

<210> 28
<211> 11
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 28

Glu Met Ala Pro Gly Ala Val His Leu Pro Gln
1 5 10

<210> 29
<211> 11
<212> PRT
<213> Artificial sequence

<220>
<223> peptide substrate

<400> 29

Glu Pro Leu Ala Gln Ala Val Arg Ser Ser Ser
1 5 10

<210> 30
<211> 11
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 30

Glu Pro Pro Val Ala Ala Ser Ser Leu Arg Asn
1 5 10

<210> 31
<211> 11
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 31

Glu Pro Gln Ile Glu Asn Val Lys Gly Thr Glu
1 5 10

<210> 32
<211> 11
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 32

Glu Ser Leu Pro Val Gln Asp Ser Ser Ser Val
1 5 10

<210> 33
<211> 11
<212> PRT
<213> artificial sequence

<220>

<223> peptide substrate

<400> 33

Glu Val His His Gln Lys Leu Val Phe Phe Ala
1 5 10

<210> 34

<211> 13

<212> PRT

<213> artificial sequence

<220>

<223> peptide substrate

<400> 34

Lys Arg Gly Val Val Asn Ala Ser Ser Arg Leu Ala Lys
1 5 10

<210> 35

<211> 9

<212> PRT

<213> artificial sequence

<220>

<223> peptide substrate

<400> 35

Lys Leu Val Leu Ala Ser Ser Ser Phe
1 5

<210> 36

<211> 13

<212> PRT

<213> artificial sequence

<220>

<223> peptide substrate

<400> 36

Lys Ser Asn Arg Leu Glu Ala Ser Ser Arg Ser Ser Pro
1 5 10

<210> 37

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> peptide substrate

<220>
<221> MISC_FEATURE
<222> (7)..(7)
<223> "x" = aminobutyric acid

<400> 37

Glu Asp Glu Met Glu Glu Xaa Ala Ser His Leu Pro Tyr
1 5 10

<210> 38
<211> 13
<212> PRT
<213> Artificial sequence

<220>
<223> peptide substrate

<400> 38

Glu Ala Gly Pro Arg Gly Met Ala Gly Gln Phe Ser His
1 . 5 10

<210> 39
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> peptide substrate

<400> 39

Lys Arg Pro Leu Gly Leu Ala Arg
1 5

<210> 40
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> peptide substrate

<400> 40

Glu Gly Tyr Tyr Ser Arg Asp Met Leu Val
1 5 10

<210> 41
<211> 11

<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 41

Glu Gln Lys Leu Asp Lys Ser Phe Ser Met Ile
1 5 10

<210> 42
<211> 12
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 42

Glu Pro Ser Ala Ala Gln Thr Ala Arg Gln His Pro
1 5 10

<210> 43
<211> 11
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 43

Glu Pro Gly Ala Gln Gly Leu Pro Gly Val Gly
1 5 10

<210> 44
<211> 8
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 44

Gly Leu Arg Thr Asn Ser Phe Ser
1 5

<210> 45
<211> 11
<212> PRT

<213> artificial sequence

<220>

<223> peptide substrate

<400> 45

Arg Gly Val Val Asn Ala Ser Ser Arg Leu Ala
1 5 10

<210> 46

<211> 10

<212> PRT

<213> artificial sequence

<220>

<223> peptide substrate

<400> 46

Lys Pro Ile Leu Phe Phe Arg Leu Gly Lys
1 5 10

<210> 47

<211> 13

<212> PRT

<213> artificial sequence

<220>

<223> peptide substrate

<400> 47

Glu Met His Thr Ala Ser Ser Leu Glu Lys Gln Ile Gly
1 5 10

<210> 48

<211> 11

<212> PRT

<213> artificial sequence

<220>

<223> peptide substrate

<400> 48

Glu Arg Phe Ala Gln Ala Gln Gln Gln Leu Pro
1 5 10

<210> 49

<211> 13

<212> PRT

<213> artificial sequence

<220>
<223> peptide substrate

<400> 49

Glu Lys Lys Glu Asn Ser Phe Glu Met Gln Gly Asp Gln
1 5 10

<210> 50
<211> 10
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 50

Leu Ala Gln Ala Val Arg Ser Ser Ser Arg
1 5 10

<210> 51
<211> 9
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 51

Glu Arg Thr Ala Ala Val Phe Arg Pro
1 5

<210> 52
<211> 8
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 52

Glu Arg Val Arg Arg Ala Leu Pro
1 5

<210> 53
<211> 9
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 53

Glu Ser Phe Pro Arg Met Phe Ser Asp
1 5

<210> 54
<211> 11
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 54

Glu Glu Tyr Leu Glu Ser Phe Leu Glu Arg Pro
1 5 10

<210> 55
<211> 12
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 55

Glu Arg Pro Lys Pro Gln Gln Phe Phe Gly Leu Met
1 5 10

<210> 56
<211> 13
<212> PRT
<213> artificial sequence

<220>
<223> peptide substrate

<400> 56

Glu His Gly Asp Gln Met Ala Gln Lys Ser Gln Ser Thr
1 5 10

<210> 57
<211> 15
<212> PRT
<213> artificial sequence

<220>

<223> peptide substrate

<400> 57

Glu Arg Asn Ile Thr Glu Gly Glu Ala Arg Gly Ser Val Ile Leu
1 5 10 15

<210> 58

<211> 10

<212> PRT

<213> artificial sequence

<220>

<223> peptide substrate

<400> 58

Glu Ala Gly Gln Arg Leu Ala Thr Ala Met
1 . 5 10

<210> 59

<211> 12

<212> PRT

<213> artificial sequence

<220>

<223> peptide substrate

<400> 59

Glu Val Gly Leu Met Gly Lys Arg Ala Leu Asn Ser
1 . 5 10

<210> 60

<211> 12

<212> PRT

<213> artificial sequence

<220>

<223> peptide substrate

<400> 60

Glu Lys Glu Asp Gly Glu Ala Arg Ala Ser Thr Ser
1 . 5 10